

②

Errors Corrected by the STIC Systems Branch

OIPE #2

Serial Number: 09/828, 498

CRF Processing Date:

Edited by:

Verified by: _____ (STIC staff)

Changed a file from non-ASCII to ASCII

Changed the margins in cases where the sequence text was "wrapped" down to the next line.

Edited a format error in the Current Application Data section, specifically: **ENTERED**

Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other _____

Added the mandatory heading and subheadings for "Current Application Data".

Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____

Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____

Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____

Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

Inserted colons after headings/subheadings. Headings edited included: _____

Deleted extra, invalid, headings used by an applicant, specifically: _____

Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file; page numbers throughout text; other invalid text, such as _____

Inserted mandatory headings, specifically: _____

Corrected an obvious error in the response, specifically: _____

Edited identifiers where upper case is used but lower case is required, or vice versa.

Corrected an error in the Number of Sequences field, specifically: _____

A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

Other: Incorrect line length error was corrected for <120> Title of Invention.

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

OIPE

RAW SEQUENCE LISTING DATE: 05/01/2001
PATENT APPLICATION: US/09/828,498 TIME: 10:58:32

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Output Set: N:\CRF3\05012001\I828498.raw

3 <110> APPLICANT: XIANG, JINHUA
4 WUNSCHMANN, SABINA
5 SCHMIDT, WARREN
6 STAPLETON, JACK T.
8 <120> TITLE OF INVENTION: FULL-LENGTH GB VIRUS C (HEPATITIS G VIRUS) RNA TRANSCRIPTS
9 ARE INFECTIOUS IN PRIMARY CD4. POSITIVE T CELLS
11 <130> FILE REFERENCE: IOWA:030US
C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/828,498
14 <141> CURRENT FILING DATE: 2001-04-05
16 <150> PRIOR APPLICATION NUMBER: 60/253,390
17 <151> PRIOR FILING DATE: 2000-11-27
19 <150> PRIOR APPLICATION NUMBER: 60/195,597
20 <151> PRIOR FILING DATE: 2000-04-06
22 <160> NUMBER OF SEQ ID NOS: 2
24 <170> SOFTWARE: PatentIn Ver. 2.1
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193 <212> TYPE: PRT

194 <213> ORGANISM: Hepatitis G virus

196 <400> SEQUENCE: 2

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 249 275 280 285
 251 Trp Gln Ser Cys Ser Cys Arg Ala Asn Gly Ser Arg Ile Pro Thr Gly
 252 290 295 300
 254 Glu Arg Val Trp Asp Arg Gly Asn Val Thr Leu Leu Cys Asp Cys Pro
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 258 325 330 335
 260 Gly Asp Pro Ile Thr His Trp Ser His Gly Gln Asn Gln Trp Pro Leu
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 263 Ser Cys Pro Gln Tyr Val Tyr Gly Ser Val Ser Val Thr Cys Val Trp
 264 355 360 365
 266 Gly Ser Val Ser Trp Phe Ala Ser Thr Gly Gly Arg Asp Ser Lys Ile
 267 370 375 380
 269 Asp Val Trp Ser Leu Val Pro Val Gly Ser Ala Ser Cys Thr Ile Ala
 270 385 390 395 400
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L:13 M:270 C: Current Application Number differs, Replaced Current Application Number